

1/23

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MAY 1 2 2003

TECH CENTER 1600/2900

Gag AF110965 BW mod

ATGGGCGCCGCGCAGCATCCTGCGCGGCGCAAGCTGGACGCCTGGGAGCGCATCCGCC TGCGCCCCGGCGGCAAGAAGTGCTACATGATGAAGCACCTGGTGTGGGCCAGCCGCGAGCT GGAGAAGTTCGCCCTGAACCCCGGCCTGCTGGAGACCAGCGAGGGCTGCAAGCAGATCATC CGCCAGCTGCACCCCGCCCTGCAGACCGGCAGCGAGGAGCTGAAGAGCCTGTTCAACACCG TGGCCACCCTGTACTGCGTGCACGAGAAGATCGAGGTCCGCGACACCAAGGAGGCCCTGGA CAAGATCGAGGAGGAGCAGAACAAGTGCCAGCAGAAGATCCAGCAGGCCGAGGCCGCCGAC AAGGGCAAGGTGAGCCAGAACTACCCCATCGTGCAGAACCTGCAGGGCCAGATGGTGCACC AGGCCATCAGCCCCCGCACCCTGAACGCCTGGGTGAAGGTGATCGAGGAGAAGGCCTTCAG CCCCGAGGTGATCCCCATGTTCACCGCCCTGAGCGAGGGCGCCACCCCCCAGGACCTGAAC ACGATGTTGAACACCGTGGGCGGCCACCAGGCCGCCATGCAGATGCTGAAGGACACCATCA ACGAGGAGGCCGCCGAGTGGGACCGCGTGCACCCCGTGCACGCCGGCCCCATCGCCCCCGG CCAGATGCGCGAGCCCCGCGGCAGCACCCGCCACCAGCACCCTGCAGGAGCAG ATCGCCTGGATGACCAGCAACCCCCCCATCCCCGTGGGCGACATCTACAAGCGGTGGATCA TCCTGGGCCTGAACAAGATCGTGCGGATGTACAGCCCCGTGAGCATCCTGGACATCAAGCA GGGCCCAAGGAGCCCTTCCGCGACTACGTGGACCGCTTCTTCAAGACCCTGCGCGCGAG CAGAGCACCCAGGAGGTGAAGAACTGGATGACCGACACCCTGCTGGTGCAGAACGCCAACC CCGACTGCAAGACCATCCTGCGCGCTCTCGGCCCCGGCGCCAGCCTGGAGGAGATGATGAC CGCCTGCCAGGCGTGGGCGCCCCAGCCACAAGGCCCGCGTGCTGGCCGAGGCGATGAGC CAGGCCAACACCAGCGTGATGATGCAGAAGAGCAACTTCAAGGGCCCCCGGCGCATCGTCA GGGCTGCTGGAAGTGCGGCAAGGAGGGCCACCAGATGAAGGACTGCACCGAGCGCCAGGCC AACTTCCTGGGCAAGATCTGGCCCAGCCACAAGGGCCGCCCCGGCAACTTCCTGCAGAGCC GCCCGAGCCCACCGCCCCCCCCGCGAGAGCTTCCGCTTCGAGGAGACCACCCCCGGCCA GAAGCAGGAGCAAGGACCGCGAGACCCTGACCAGCCTGAAGAGCCTGTTCGGCAACGAC CCCCTGAGCCAGTAA



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ATGGGCGCCCGCGCATCCTGCGCGCGAGAAGCTGGACAAGTGGGAGAAGATCCGCC TGCGCCCGGCGGCAAGAAGCACTACATGCTGAAGCACCTGGTGTGGGCCAGCCGCGAGCT GGAGGGCTTCGCCCTGAACCCCGGCCTGCTGGAGACCGCCGAGGGCTGCAAGCAGATCATG AAGCAGCTGCAGCCCGCCCTGCAGACCGGCAGCGAGCGTGCAGCCTGTACAACACCG TGGCCACCCTGTACTGCGTGCACGCCGGCATCGAGGTCCGCGACACCAAGGAGGCCCTGGA CAAGATCGAGGAGGAGCAGAACAAGTCCCAGCAGAAGACCCAGCAGGCCAAGGAGGCCGAC GGCAAGGTGAGCCAGAACTACCCCATCGTGCAGAACCTGCAGGGCCAGATGGTGCACCAGG CCATCAGCCCCGCACCCTGAACGCCTGGGTGAAGGTGATCGAGGAGAAGGCCTTCAGCCC CGAGGTGATCCCCATGTTCACCGCCCTGAGCGAGGGCGCCACCCCCAGGACCTGAACACG ATGTTGAACACCGTGGGCGGCCACCAGGCCGCCATGCAGATGCTGAAGGACACCATCAACG GATGCGCGACCCCCGCGGCAGCACCATCGCCGGCGCCACCAGCACCCTGCAGGAGCAGATC GCCTGGATGACCAGCAACCCCCCGTGCCCGTGGGCGACATCTACAAGCGGTGGATCATCC TGGGCCTGAACAAGATCGTGCGGATGTACAGCCCCGTGAGCATCCTGGACATCCGCCAGGG CCCCAAGGAGCCCTTCCGCGACTACGTGGACCGCTTCTTCAAGACCCTGCGCGCGAGCAG GCCACCCAGGACGTGAAGAACTGGATGACCGAGACCCTGCTGGTGCAGAACGCCAACCCCG ACTGCAAGACCATCCTGCGCGCTCTCGGCCCCCGGCGCCACCCTGGAGGAGATGATGACCGC $\tt CTGCCAGGGCGTGGGGGGGGGCCCACAAGGCCCGCGTGCTGGCCGAGGCGATGAGCCAG$ GCCAACAGCGTGAACATCATGATGCAGAAGAGCAACTTCAAGGGCCCCCGGCGCAACGTCA GGGCTGCTGGAAGTGCGGCAAGGAGGGCCACCAGATGAAGGACTGCACCGAGCGCCAGGCC AACTTCCTGGGCAAGATCTGGCCCAGCCACAAGGGCCGCCCCGGCAACTTCCTGCAGAACC GGAGACCACCCCCCCCAAGCAGGAGCCCAAGGACCGCGAGCCCTACCGCGAGCCCCTG ACCGCCCTGCGCAGCCTGTTCGGCAGCCGCCCCTGAGCCAGTAA



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--> signal peptide (1-81)
ATGCGCGTGATGGGCATCCTGAAGAACTACCAGCAGTGGTGGATGTGGGGCATCCTGGGCTTCTGGATGCTGATCA V--> gp120/140/160 (82)
TCAGCAGCGTGGTGGCAACCTGTGGGTGACCACCCT GACCCCAACCCCAGGAGATCGTGCTGGAGAACGTGACCGAGAACTTCAACATGTGGAAGAACGACATGGTGGACC AGATGCACGAGGACATCATCAGCCTGTGGGACCAGAGCCTGAAGCCCTGCGTGAAGCTGACCCCCTGTGCGTGAC CCTGAAGTGCCGCAACGTGAACGCCACCAACAACATCAACAGCATGATCGACAACAGCAACAAGGGCGAGATGAAG AACTGCAGCTTCAACGTGACCACCGAGCTGCGCGACCGCAAGCAGGAGGTGCACGCCCTGTTCTACCGCCTGGACG TGGTGCCCCTGCAGGGCAACAACAGCAACGAGTACCGCCTGATCAACTGCAACACCAGCGCCATCACCCAGGCCTG CCCCAAGGTGAGCTTCGACCCCATCCCATCCACTACTGCACCCCGGCGGCTACGCCATCCTGAAGTGCAACAAC CAGACCTTCAACGGCACCGGCCCTGCAACAACGTGAGCAGCGTGCAGTGCGCCCCACGGCATCAAGCCCGTGGTGA GCACCCAGCTGCTGAACGGCAGCCTGGCCAAGGGCGAGATCATCATCCGCAGCGAGAACCTGGCCAACAACGC GTGCGCATCGGCCCGGCCAGACCTTCTACGCCACCGGCGAGATCATCGGCGACATCCGCCAGGCCTACTGCATCA TCAACAAGACCGAGTGGAACAGCACCCTGCAGGGCGTGAGCAAGAAGCTGGAGGAGCACTTCAGCAAGAAGGCCAT TGCGACACCAGCCAGCTGTTCAACAGCACCTACAGCCCCAGCTTCAACGGCACCGAGAACAAGCTGAACGGCACCA CGCCGGCAACCTGACCTGCGAGAGCAACATCACCGGCCTGCTGACCCGCGACGGCGGCAAGACCCGACCAAC GACACCGAGATCTTCCGCCCGGCGGCGGCGACATGCGCGACAACTGGCGCAACGAGCTGTACAAGTACAAGGTGG TGGAGATCAAGCCCCTGGGCGTGGCCCCCACCGAGGCCAAGCGCCGCGTGGTGGAGCGCGAGAAGCGCGCGTGGG CATCGGCGCCGTGTTCCTGGGCTTCCTGGGCGCCGCCGCCAGCATCACCCTGACCGTG CAGGCCCGCCTGCTGAGCGGCATCGTGCAGCAGCAGCAACAACCTGCTGCGCGCCATCGAGGCCCAGCAGCACC TGCTGCAGCTGACCGTGTGGGGCATCAAGCAGCTGCAGACCCGCATCCTGGCCGTGGAGCGCTACCTGAAGGACCA GCAGCTGCTGGGCATCTGGGGCTGCAGCGGCAAGCTGATCTGCACCACCGCCGTGCCCTGGAACAGCAGCTGGAGC AACCGCAGCCACGACGAGATCTGGGACAACATGACCTGGATGCAGTGGGACCGCGAGATCAACAACTACACCGACA CCATCTACCGCCTGCTGGAGGAGAGCCAGAACCAGCAGGAGAAGAACGAGAAGGACCTGCTGGCCCTGGACAGCTG GCAGAACCTGTGGAACTGGTTCAGCATCACCAACTGGCTGTGGTACATCAAGATCTTCATCATGATCGTGGGCGGC CTGATCGGCCTGCGCATCATCTTCGCCGTGCTGAGCATCGTGAACCGCGTGCGCCAGGGCTACAGCCCCCTGCCCT CGGCCGCAGCATCCGCCTGGTGAGCGGCTTCCTGGCCCTGGCCTGGGACGACCTGCGCAGCCTGTTCAGC TACCACCGCCTGCGCGACTTCATCCTGATCGCCGCCCGCGTGCTGGAGCTGCTGGGCCAGCGCGGCTGGGAGGCCC TGAAGTACCTGGGCAGCCTGGTGCAGTACTGGGGCCTGGAGCTGAAGAAGAGGCGCCATCAGCCTGCTGGACACCAT $\tt CGCCATCGCCGAGGGCACCGACCGCATCATCGAGTTCATCCAGCGCATCTGCCGCGCCATCCGCAACATC$ GP160, GP41(2547)<--\
CCCCGCCGCATCCGCCAGGGCTTCGAGGCCGCCCTGCAGTAA

MAY 0 8 2003

Polynucleotides Encoding Antigenic HIV Type C Polypeptides, Polypeptides and Uses Thereof

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--> signal peptide (1-72)
ATGCGCGTGCGCGCATCCTGCGCAGCTGGCAGCAGCTGGATCTGGGGCATCCTGGGCTTCTGGATCTGCAGCG gp120/140/160 (72)
GCCTGGGCAACCTGTGACCGTGTACGACGGCGTGCCCGTGTGGCGCGAGGCCACCACCCTGTTCTGCGC CCCCAGGAGATCGAGCTGGACAACGTGACCGAGAACTTCAACATGTGGAAGAACGACATGGTGGACCAGATGCACG AGGACATCATCAGCCTGTGGGACCAGAGCCTGAAGCCCCGCGTGAAGCTGACCCCCCTGTGCGTGACCCTGAAGTG CACCAACTACAGCACCAACTACAGCAACACCATGAACGCCACCAGCTACAACAACAACACCACCGAGGAGATCAAG AACTGCACCTTCAACATGACCACCGAGCTGCGCGACAAGAAGCAGCAGGTGTACGCCCTGTTCTACAAGCTGGACA TCGTGCCCCTGAACAGCAACAGCGGGGTACCGCCTGATCAACTGCAACACCAGCGCCATCACCCAGGCCTGCCC CAAGGTGAGCTTCGACCCCATCCCATCCACTACTGCGCCCCGCCGGCTACGCCATCCTGAAGTGCAAGAACAAC ACCAGCAACGGCACCGGCCCCTGCCAGAACGTGAGCACCGTGCAGTGCACCCACGGCATCAAGCCCGTGGTGAGCA CTACACCATCATCGTGCACCTGAACGACAGCGTGGAGATCGTGTGCACCCGCCCAACAACAACACCCGCAAGGGC ATCCGCATCGGCCCGGCCAGACCTTCTACGCCACCGAGAACATCATCGGCGACATCCGCCAGGCCCACTGCAACA TCAGCGCCGGCGAGCTGGAACAAGGCCGTGCAGCGCGTGAGCGCCAAGCTGCGCGAGCACTTCCCCAACAAGACCAT TGCAACACCAGCAAGCTGTTCAACAGCAGCTACAACGGCACCAGCTACCGCGGCACCGAGAGCAACAGCAGCATCA TCACCCTGCCCTGCCGCATCAAGCAGATCATCGACATGTGGCAGAAGGTGGGCCGCCCCCATCTACGCCCCCCCAT CGAGGGCAACATCACCTGCAGCAGCAGCATCACCGGCCTGCTGCTGGCCGGCGGCGGGCCTGGACAACATCACC ACCGAGATCTTCCGCCCCAGGGCGGCGACATGAAGGACAACTGGCGCAACGAGCTGTACAAGTACAAGGTGGTGG AGATCAAGCCCCTGGGCGTGGCCCCCACCGAGGCCAAGCGCCGCGTGGTGGAGCGCGAGAAGCGCGCGTGGGCAT CGGCGCCGTGATCTTCGGCTTCCTGGGCGCCGCCGGCAGCAACATGGGCGCCGCCAGCATCACCCTGACCGCCCAG TGCAGCTGACCGTGTGGGGCATCAAGCAGCTGCAGGCCCGCGTGCTGGCCATCGAGCGCTACCTGAAGGACCAGCA GCTGCTGGGCATCTGGGGCTGCAGCGGCAAGCTGATCTGCACCACCGTGCCCTGGAACAGCAGCTGGAGCAAC AAGACCCAGGGCGAGATCTGGGAGAACATGACCTGGATGCAGTGGGACAAGGAGATCAGCAACTACACCGGCATCA TCTACCGCCTGCTGGAGGAGGCCAGAACCAGCAGGAGCAGAACGAGAAGGACCTGCTGGCCCTGGACAGCCGCAA GP140(2022)<--\/
CAACCTGTGGAGCTGGTTCAACATCAGCAACTGGCTGGTACATCAAGATCTTCATCATGATCGTGGGCGGCCTG ATCGGCCTGCGCATCATCTTCGCCGTGCTGAGCATCGTGAACCGCGTGCGCCAGGGCTACAGCCCCCTGAGCTTCC CACCGCCTGCGCGACCTGATCCTGGTGACCGCCGCGTGGTGGAGCTGCTGGGCCGCAGCAGCCCCCGCGGCCTGC AGCGCGGCTGGGAGCCCTGAAGTACCTGGGCAGCCTGGTGCAGTACTGGGGCCTGGAGCTGAAGAAGAGCGCCAC gp160, gp41(2565)<--\
CGCGCCTTCTGCAACATCCCCGCCGCGTGCGCCAGGGCTTCGAGGCCGCCCTGCAGTAA



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Gag_AF110965_BW_opt ATGGGCGCCGCGCCAGCATCCTGCGCGGCGCAAGCTGGACGCCTGGGAGCGCATCCGCCTGCGCCCCGG CGGCAAGAAGTGCTACATGATGAAGCACCTGGTGTGGGCCAGCCGCGAGCTGGAGAAGTTCGCCCTGAACC CCGGCCTGCTGGAGACCAGCGAGGGCTGCAAGCAGATCATCCGCCAGCTGCACCCCGCCCTGCAGACCGGC AGCGAGGAGCTGAAGAGCCTGTTCAACACCGTGGCCACCCTGTACTGCGTGCACGAGAAGATCGAGGTGCG CGACACCAAGGAGGCCCTGGACAAGATCGAGGAGGAGCAGAACAAGAGCCAGCAGAAGATCCAGCAGGCCG CAGGCCATCAGCCCCGCACCCTGAACGCCTGGGTGAAGGTGATCGAGGAGAAGGCCTTCAGCCCCGAGGT GATCCCCATGTTCACCGCCCTGAGCGAGGGCGCCACCCCCAGGACCTGAACACCATGCTGAACACCGTGG GCGGCCACCAGGCCGCCATGCAGATGCTGAAGGACACCATCAACGAGGAGGCCGCCGAGTGGGACCGCGTG CACCCCGTGCACGCCGGCCCATCGCCCCGGGCCAGATGCGCGAGCCCCGGGCAGCACATCGCCGGCAC CACCAGCACCCTGCAGGAGCAGATCGCCTGGATGACCAGCAACCCCCCATCCCCGTGGGCGACATCTACA AGCGCTGGATCATCCTGGGCCTGAACAAGATCGTGCGCATGTACAGCCCCGTGAGCATCCTGGACATCAAG CCAGGAGGTGAAGAACTGGATGACCGACACCCTGCTGGTGCAGAACGCCAACCCCGACTGCAAGACCATCC TGCGCGCCTGGGCCCCGGCGCCAGCCTGGAGGAGATGATGACCGCCTGCCAGGGCGTGGGCGGCCCCAGC $\mathsf{CACAAGGCCCGCGTGCTGGCCGAGGGC}$ $\mathsf{CACAGGCCAGCGTGATGATGCAGAAGAGCAACTT}$ CAAGGGCCCCCGCGCGCATCGTGAAGTGCTTCAACTGCGGCAAGGAGGGCCACATCGCCCGCAACTGCCGCG CCCCCGCAAGAAGGGCTGCTGGAAGTGCGGCAAGGAGGCCCACCAGATGAAGGACTGCACCGAGCGCCAG GCCAACTTCCTGGGCAAGATCTGGCCCAGCCACAAGGGCCGCCCCGGCAACTTCCTGCAGAGCCGCCCCGA GCCCACCGCCCCCCCGGCGAGAGCTTCCGCTTCGAGGAGACCACCCCCGGCCAGAAGCAGGAGAGCAAGG ACCGCGAGACCCTGACCAGCCTGAAGAGCCTGTTCGGCAACGACCCCCTGAGCCAGTAA

MAY 0 8 2000

Polynucleotides Encoding Antigenic HIV Type C Polypeptides, Polypeptides and Uses Thereof

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Gag_AF110967_BW_opt ATGGGCGCCGCGCAGCATCCTGCGCGGGGAGAAGCTGGACAAGTGGGAGAAGATCCGCCTGCGCCCCGG CGGCAAGAAGCACTACATGCTGAAGCACCTGGTGTGGGGCCAGCCGCGAGCTGGAGGGCTTCGCCCTGAACC ACCGAGGAGCTGCGCAGCCTGTACAACACCGTGGCCACCCTGTACTGCGTGCACGCCGGCATCGAGGTGCG ${\tt AGGAGGCCGACGGCAAGGTGAGCCAGAACTACCCCATCGTGCAGAACCTGCAGGGCCAGATGGTGCACCAG}$ GCCATCAGCCCCGCACCCTGAACGCCTGGGTGAAGGTGATCGAGGAGAAGGCCTTCAGCCCCGAGGTGAT CCCCATGTTCACCGCCCTGAGCGAGGGCGCCACCCCCAGGACCTGAACACCATGGTGAACACCGTGGGCG CAGCACCCTGCAGGAGCAGATCGCCTGGATGACCAGCAACCCCCCGTGCCCGTGGGCGACATCTACAAGC GCTGGATCATCCTGGGCCTGAACAAGATCGTGCGCATGTACAGCCCCGTGAGCATCCTGGACATCCGCCAG GGACGTGAAGAACTGGATGACCGAGACCCTGCTGGTGCAGAACGCCAACCCCGACTGCAAGACCATCCTGC GCGCCTGGGCCCCCGGCCCCCTGGAGGAGATGATGACCGCCTGCCAGGGCGTGGGCGGCCCCGGCCAC AAGGCCCGCGTGCTGGCCGAGGCCATGAGCCAGCCCAACAGCGTGAACATCATGATGCAGAAGAGCAACTT CAAGGGCCCCCCCCCCAACGTGAAGTGCTTCAACTGCGGCAAGGAGGGCCACATCGCCAAGAACTGCCGCG CCCCCGCAAGAAGGCTGCTGGAAGTGCGGCAAGGAGGCCCACCAGATGAAGGACTGCACCGAGCGCCAG GCCAACTTCCTGGGCAAGATCTGGCCCAGCCACAAGGGCCGCCCGGCAACTTCCTGCAGAACCGCAGCGA CCAAGCAGGAGCCCAAGGACCGCGAGCCCTACCGCGAGCCCTGACCGCCCTGCGCAGCCTGTTCGGCAGC GGCCCCTGAGCCAGTAA



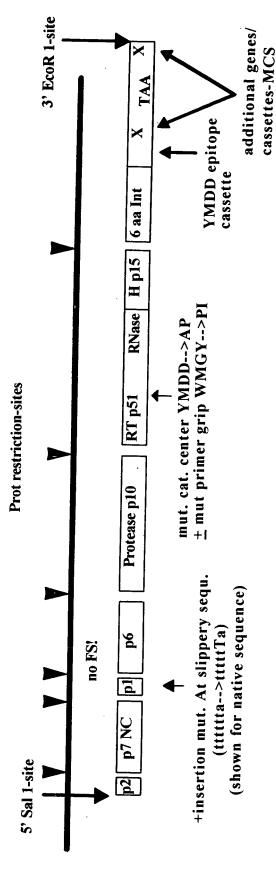


FIG. 7



8/23

PR975(+) (SEQ ID NO:30)

GTCGACGCCACCATGGCCGAGGCCATGAGCCAGGCCACCAGCGCCAACATCCTGAT GCAGCGCAGCAACTTCAAGGGCCCCAAGCGCATCATCAAGTGCTTCAACTGCGGCAA GGAGGGCCACATCGCCGCAACTGCCGCGCCCCCCGCAAGAAGGGCTGCTGGAAGT GCGGCAAGGAGGCCACCAGATGAAGGACTGCACCGAGCGCCAGGCCAACTTCTTC CGCGAGGACCTGGCCTTCCCCCAGGGCAAGGCCCGCGAGTTCCCCAGCGAGCAGAA CCGCGCCAACAGCCCACCAGCCGCGAGCTGCAGGTGCGCGGCGACAACCCCCGCA GCGAGGCCGGCCCGAGCGCCAGGGCACCCTGAACTTCCCCCAGATCACCCTGTGGC AGCGCCCCTGGTGAGCATCAAGGTGGGCGGCCAGATCAAGGAGGCCCTGCTGGAC ACCGGCGCCGACGACACCGTGCTGGAGGAGATGAGCCTGCCCGGCAAGTGGAAGCC CAAGATGATCGGCGGCATCGGCGGCTTCATCAAGGTGCGCCAGTACGACCAGATCCT GATCGAGATCTGCGGCAAGAAGGCCATCGGCACCGTGCTGATCGGCCCCACCCCCGT GAACATCATCGGCCGCAACATGCTGACCCAGCTGGGCTGCACCCTGAACTTCCCCAT CAGCCCCATCGAGACCGTGCCCGTGAAGCTGAAGCCCGGCATGGACGGCCCCAAGG TGAAGCAGTGGCCCCTGACCGAGGAGAAGATCAAGGCCCTGACCGCCATCTGCGAG GAGATGGAGAAGGGGCAAGATCACCAAGATCGGCCCCGAGAACCCCTACAACAC CCCCGTGTTCGCCATCAAGAAGAAGGACAGCACCAAGTGGCGCAAGCTGGTGGACT TCCGCGAGCTGAACAAGCGCACCCAGGACTTCTGGGAGGTGCAGCTGGGCATCCCCC ACCCCGCCGGCCTGAAGAAGAAGAAGAGCGTGACCGTGCTGGACGTGGGCGACGCC TACTTCAGCGTGCCCCTGGACGAGGACTTCCGCAAGTACACCGCCTTCACCATCCCC AGCATCAACAACGAGACCCCCGGCATCCGCTACCAGTACAACGTGCTGCCCCAGGGC TGGAAGGCCAGCATCTTCCAGAGCAGCATGACCAAGATCCTGGAGCCCTTC CGCGCCCGCAACCCCGAGATCGTGATCTACCAGTACATGGACGACCTGTACGTGGGC AGCGACCTGGAGATCGGCCAGCACCGCGCCAAGATCGAGGAGCTGCGCAAGCACCT GCTGCGCTGGGGCTTCACCACCCCGACAAGAAGCACCAGAAGGAGCCCCCCTTCCT GTGGATGGGCTACGAGCTGCACCCCGACAAGTGGACCGTGCAGCCCATCGAGCTGCC CGAGAAGGAGAGCTGGACCGTGAACGACATCCAGAAGCTGGTGGGCAAGCTGAACT GGGCCAGCCAGATCTACCCCGGCATCAAGGTGCGCCAGCTGTGCAAGCTGCTGCGCG GCGCCAAGGCCCTGACCGACATCGTGCCCCTGACCGAGGGGGGCCGAGCTGGAGCTG GCCGAGAACCGCGAGATCCTGCGCGAGCCCGTGCACGGCGTGTACTACGACCCCAG CAAGGACCTGGTGGCCGAGATCCAGAAGCAGGGCCACGACCAGTGGACCTACCAGA TCTACCAGGAGCCCTTCAAGAACCTGAAGACCGGCAAGTACGCCAAGATGCGCACC GCCCACACCAACGACGTGAAGCAGCTGACCGAGGCCGTGCAGAAGATCGCCATGGA GAGCATCGTGATCTGGGGCAAGACCCCCAAGTTCCGCCTGCCCATCCAGAAGGAGAC CTGGGAGACCTGGTGGACCGACTACTGGCAGGCCACCTGGATCCCCGAGTGGGAGTT CGTGAACACCCCCCCCCTGGTGAAGCTGTGGTACCAGCTGGAGAAGGAGCCCATCAT CGGCGCGAGACCTTCTACGTGGACGGCGCCCCAACCGCGAGACCAAGATCGGCA AGGCCGGCTACGTGACCGACCGGGGCCGGCAGAAGATCGTGAGCCTGACCGAGACC ACCAACCAGAAGACCGAGCTGCAGGCCATCCAGCTGGCCCTGCAGGACAGCGGCAG CGAGGTGAACATCGTGACCGACAGCCAGTACGCCCTGGGCATCATCCAGGCCCAGCC CGACAAGAGCGAGCGAGCTGGTGAACCAGATCATCGAGCAGCTGATCAAGAAGG AGAAGGTGTACCTGAGCTGGGTGCCCGCCCACAAGGGCATCGGCGGCAACGAGCAG ATCGACAAGCTGGTGAGCAAGGGCATCCGCAAGGTGCTGTTCCTGGACGGCATCGAT GGCGGCATCGTGATCTACCAGTACATGGACGACCTGTACGTGGGCAGCGGCGGCCCT AGGATCGATTAAAAGCTTCCCGGGGCTAGCACCGGTGAATTC



9/23

PR975YM (SEQ ID NO:31)

GTCGACGCCACCATGGCCGAGGCCATGAGCCAGGCCACCAGCGCCAACATCCTGAT GCAGCGCAGCAACTTCAAGGGCCCCAAGCGCATCATCAAGTGCTTCAACTGCGGCAA GGAGGCCACATCGCCCGCAACTGCCGCGCCCCCCGCAAGAAGGGCTGCTGGAAGT GCGGCAAGGAGGCCACCAGATGAAGGACTGCACCGAGCGCCAGGCCAACTTCTTC CGCGAGGACCTGGCCTTCCCCCAGGGCAAGGCCCGCGAGTTCCCCAGCGAGCAGAA CCGCGCCAACAGCCCACCAGCCGCGAGCTGCAGGTGCGCGGCGACAACCCCCGCA GCGAGGCCGCGCGAGCGCCAGGGCACCCTGAACTTCCCCCAGATCACCCTGTGGC AGCGCCCCTGGTGAGCATCAAGGTGGGCGGCCAGATCAAGGAGGCCCTGCTGGAC ACCGGCGCCGACGACACCGTGCTGGAGGAGATGAGCCTGCCCGGCAAGTGGAAGCC CAAGATGATCGGCGGCATCGGCGGCTTCATCAAGGTGCGCCAGTACGACCAGATCCT GATCGAGATCTGCGGCAAGAAGGCCATCGGCACCGTGCTGATCGGCCCCACCCCCGT GAACATCATCGGCCGCAACATGCTGACCCAGCTGGGCTGCACCCTGAACTTCCCCAT CAGCCCCATCGAGACCGTGCCCGTGAAGCTGAAGCCCGGCATGGACGGCCCCAAGG TGAAGCAGTGGCCCCTGACCGAGGAGAAGATCAAGGCCCTGACCGCCATCTGCGAG GAGATGGAGAAGGGGCAAGATCACCAAGATCGGCCCCGAGAACCCCTACAACAC CCCCGTGTTCGCCATCAAGAAGAAGAAGACAGCACCAAGTGGCGCAAGCTGGTGGACT TCCGCGAGCTGAACAAGCGCACCCAGGACTTCTGGGAGGTGCAGCTGGGCATCCCCC ACCCCGCCGGCCTGAAGAAGAAGAAGAGCGTGACCGTGCTGGACGTGGGCGACGCC TACTTCAGCGTGCCCCTGGACGAGGACTTCCGCAAGTACACCGCCTTCACCATCCCC AGCATCAACAACGAGACCCCCGGCATCCGCTACCAGTACAACGTGCTGCCCCAGGGC TGGAAGGCAGCCCCAGCATCTTCCAGAGCAGCATGACCAAGATCCTGGAGCCCTTC CGCGCCCGCAACCCCGAGATCGTGATCTACCAGGCCCCCCTGTACGTGGGCAGCGAC CTGGAGATCGGCCAGCACCGCGCCAAGATCGAGGAGCTGCGCAAGCACCTGCTGCG CTGGGGCTTCACCACCCCGACAAGAAGCACCAGAAGGAGCCCCCCTTCCTGTGGAT GGGCTACGAGCTGCACCCCGACAAGTGGACCGTGCAGCCCATCGAGCTGCCCGAGA AGGAGAGCTGGACCGTGAACGACATCCAGAAGCTGGTGGGCCAAGCTGAACTGGGCC GAACCGCGAGATCCTGCGCGAGCCCGTGCACGGCGTGTACTACGACCCCAGCAAGG ACCTGGTGGCCGAGATCCAGAAGCAGGGCCACGACCAGTGGACCTACCAGATCTAC CAGGAGCCCTTCAAGAACCTGAAGACCGGCAAGTACGCCAAGATGCGCACCGCCCA CACCAACGACGTGAAGCAGCTGACCGAGGCCGTGCAGAAGATCGCCATGGAGAGCA TCGTGATCTGGGGCAAGACCCCCAAGTTCCGCCTGCCCATCCAGAAGGAGACCTGGG AGACCTGGTGGACCGACTACTGGCAGGCCACCTGGATCCCCGAGTGGGAGTTCGTGA ACACCCCCCCCTGGTGAAGCTGTGGTACCAGCTGGAGAAGGAGCCCATCATCGGCG CCGAGACCTTCTACGTGGACGCCCCCCCCAACCGCGAGACCAAGATCGGCAAGGCC GGCTACGTGACCGACCGGGCCGGCAGAAGATCGTGAGCCTGACCGAGACCACCAA TGAACATCGTGACCGACAGCCAGTACGCCCTGGGCATCATCCAGGCCCAGCCCGACA AGAGCGAGAGCGAGCTGGTGAACCAGATCATCGAGCAGCTGATCAAGAAGGAGAAG GTGTACCTGAGCTGGGCCCCCCCCCCACAAGGGCATCGGCGCAACGAGCAGATCGA CAAGCTGGTGAGCAAGGGCATCCGCAAGGTGCTGTTCCTGGACGCATCGATGGCG GCATCGTGATCTACCAGTACATGGACGACCTGTACGTGGGCAGCGGCGCCCTAGGA TCGATTAAAAGCTTCCCGGGGCTAGCACCGGTGAATTC



10/23

PR975YMWM (SEQ ID NO:32)

GTCGACGCCACCATGGCCGAGGCCATGAGCCAGGCCACCAGCGCCAACATCCTGAT GCAGCGCAGCAACTTCAAGGGCCCCAAGCGCATCATCAAGTGCTTCAACTGCGGCAA GGAGGGCCACATCGCCGCAACTGCCGCGCCCCCCGCAAGAAGGGCTGCTGGAAGT GCGGCAAGGAGGCCACCAGATGAAGGACTGCACCGAGCGCCAGGCCAACTTCTTC CGCGAGGACCTGGCCTTCCCCCAGGGCAAGGCCCGCGAGTTCCCCAGCGAGCAGAA CCGCGCCAACAGCCCACCAGCCGCGAGCTGCAGGTGCGCGGCGACAACCCCCGCA GCGAGGCCGGCGCGAGCGCCAGGGCACCCTGAACTTCCCCCAGATCACCCTGTGGC AGCGCCCCTGGTGAGCATCAAGGTGGGCGGCCAGATCAAGGAGGCCCTGCTGGAC ACCGGCGCCGACGACACCGTGCTGGAGGAGATGAGCCTGCCCGGCAAGTGGAAGCC CAAGATGATCGGCGGCATCGGCGGCTTCATCAAGGTGCGCCAGTACGACCAGATCCT GATCGAGATCTGCGGCAAGAAGGCCATCGGCACCGTGCTGATCGGCCCCACCCCCGT GAACATCATCGGCCGCAACATGCTGACCCAGCTGGGCTGCACCCTGAACTTCCCCAT CAGCCCCATCGAGACCGTGCCCGTGAAGCTGAAGCCCGGCATGGACGGCCCCAAGG TGAAGCAGTGGCCCCTGACCGAGGAGAAGATCAAGGCCCTGACCGCCATCTGCGAG GAGATGGAGAAGGGGCAAGATCACCAAGATCGGCCCCGAGAACCCCTACAACAC CCCCGTGTTCGCCATCAAGAAGAAGGACAGCACCAAGTGGCGCAAGCTGGTGGACT TCCGCGAGCTGAACAAGCGCACCCAGGACTTCTGGGAGGTGCAGCTGGGCATCCCCC ACCCCGCCGGCCTGAAGAAGAAGAAGAGCGTGACCGTGCTGGACGTGGGCGACGCC TACTTCAGCGTGCCCCTGGACGAGGACTTCCGCAAGTACACCGCCTTCACCATCCCC AGCATCAACAACGAGACCCCCGGCATCCGCTACCAGTACAACGTGCTGCCCCAGGGC TGGAAGGCCAGCCCCAGCATCTTCCAGAGCAGCATGACCAAGATCCTGGAGCCCTTC CGCGCCCGCAACCCCGAGATCGTGATCTACCAGGCCCCCCTGTACGTGGGCAGCGAC CTGGAGATCGGCCAGCACCGCGCCAAGATCGAGGAGCTGCGCAAGCACCTGCTGCG CTGGGGCTTCACCACCCCGACAAGAAGCACCAGAAGGAGCCCCCCTTCCTGCCCAT CGAGCTGCACCCCGACAAGTGGACCGTGCAGCCCATCGAGCTGCCCGAGAAGGAGA CGAGATCCTGCGCGAGCCCGTGCACGGCGTGTACTACGACCCCAGCAAGGACCTGGT GGCCGAGATCCAGAAGCAGGGCCACGACCAGTGGACCTACCAGATCTACCAGGAGC CCTTCAAGAACCTGAAGACCGGCAAGTACGCCAAGATGCGCACCCCACACCAAC GACGTGAAGCAGCTGACCGAGGCCGTGCAGAAGATCGCCATGGAGAGCATCGTGAT CTGGGGCAAGACCCCAAGTTCCGCCTGCCCATCCAGAAGGAGACCTGGGAGACCT GGTGGACCGACTACTGGCAGGCCACCTGGATCCCCGAGTGGGAGTTCGTGAACACCC CCCCCTGGTGAAGCTGTGGTACCAGCTGGAGAAGGAGCCCATCATCGGCGCCGAG ACCTTCTACGTGGACGGCGCCCAACCGCGAGACCAAGATCGGCAAGGCCGGCTA CGTGACCGACCGGGCCGGCAGAGATCGTGAGCCTGACCGAGACCACCAGA AGACCGAGCTGCAGGCCATCCAGCTGGCCCTGCAGGACAGCGGCAGCGAGGTGAAC ATCGTGACCGACAGCCAGTACGCCCTGGGCATCATCCAGGCCCAGCCCGACAAGAG CGAGAGCGAGCTGGTGAACCAGATCATCGAGCAGCTGATCAAGAAGGAGAAGGTGT ACCTGAGCTGGGTGCCCGCCCACAAGGGCATCGGCGGCAACGAGCAGATCGACAAG CTGGTGAGCAAGGCATCCGCAAGGTGCTGTTCCTGGACGGCATCGATGGCGGCATC GTGATCTACCAGTACATGGACGACCTGTACGTGGGCAGCGGCGGCCCTAGGATCGAT TAAAAGCTTCCCGGGGCTAGCACCGGTGAATTC



11/23

8_5_ZA (SEQ ID NO:33)

1 TGGAAGGGTT AATTTACTCC AAGAAAAGGC AAGAAATCCT TGATTTGTGG GTCTATCACA 61 CACAAGGCTT CTTCCCTGAT TGGCAAAACT ACACACCGGG GCCAGGGGTC AGATATCCAC 121 TGACCTTTGG ATGGTGCTAC AAGCTAGTGC CAGTTGACCC AGGGGAGGTG GAAGAGGCCA 181 ACGGAGGAGA AGACAACTGT TTGCTACACC CTATGAGCCA ACATGGAGCA GAGGATGAAG 241 ATAGAGAAGT ATTAAAGTGG AAGTTTGACA GCCTCCTAGC ACGCAGACAC ATGGCCCGCG 301 AGCTACATCC GGAGTATTAC AAAGACTGCT GACACAGAAG GGACTTTCCG CCTGGGACTT 361 TCCACTGGGG CGTTCCGGGA GGTGTGGTCT GGGCGGGACT TGGGAGTGGT CAACCCTCAG 421 ATGCTGCATA TAAGCAGCTG CTTTTCGCCT GTACTGGGTC TCTCTCGGTA GACCAGATCT 481 GAGCCTGGGA GCCCTCTGGC TATCTAGGGA ACCCACTGCT TAAGCCTCAA TAAAGCTTGC 541 CTTGAGTGCT TTAAGTAGTG TGTGCCCATC TGTTGTGTGA CTCTGGTAAC TAGAGATCCC 601 TCAGACCCTT TGTGGTAGTG TGGAAAATCT CTAGCAGTGG CGCCCGAACA GGGACCAGAA 661 AGTGAAAGTG AGACCAGAGG AGATCTCTCG ACGCAGGACT CGGCTTGCTG AAGTGCACAC 721 GGCAAGAGGC GAGAGGGCG GCTGGTGAGT ACGCCAATTT TACTTGACTA GCGGAGGCTA 781 GAAGGAGAG GATGGGTGCG AGAGCGTCAA TATTAAGCGG CGGAAAATTA GATAAATGGG 841 AAAGAATTAG GTTAAGGCCA GGGGGAAAGA AACATTATAT GTTAAAACAT CTAGTATGGG 901 CAAGCAGGGA GCTGGAAAGA TTTGCACTTA ACCCTGGCCT GTTAGAAACA TCAGAAGGCT 961 GTAAACAAAT AATAAAACAG CTACAACCAG CTCTTCAGAC AGGAACAGAG GAACTTAGAT 1021 CATTATTCAA CACAGTAGCA ACTCTCTATT GTGTACATAA AGGGATAGAG GTACGAGACA 1141 AGGCAAAAGC AGCTGACGAA AAGGTCAGTC AAAATTATCC TATAGTACAG AATGCCCAAG 1201 GGCAAATGGT ACACCAAGCT ATATCACCTA GAACATTGAA TGCATGGATA AAAGTAATAG 1261 AGGAAAAGGC TTTCAATCCA GAGGAAATAC CCATGTTTAC AGCATTATCA GAAGGAGCCA 1321 CCCCACAAGA TTTAAACACA ATGTTAAATA CAGTGGGGGG ACATCAAGCA GCCATGCAAA 1381 TGTTAAAAGA TACCATCAAT GAGGAGGCTG CAGAATGGGA TAGGACACAT CCAGTACATG 1441 CAGGGCCTGT TGCACCAGGC CAGATGAGAG AACCAAGGGG AAGTGACATA GCAGGAACTA 1501 CTAGTACCCT TCAGGAACAA ATAGCATGGA TGACAAGTAA TCCACCTATT CCAGTAGAAG 1561 ACATCTATAA AAGATGGATA ATTCTGGGGT TAAATAAAAT AGTAAGAATG TATAGCCCTG 1621 TTAGCATTTT GGACATAAAA CAAGGGCCAA AAGAACCCTT TAGAGACTAT GTAGACCGGT 1681 TCTTTAAAAC CTTAAGAGCT GAACAAGCTA CACAAGATGT AAAGAATTGG ATGACAGACA 1741 CCTTGTTGGT CCAAAATGCG AACCCAGATT GTAAGACCAT TTTAAGAGCA TTAGGACCAG 1801 GGGCCTCATT AGAAGAAATG ATGACAGCAT GTCAGGGAGT GGGAGGACCT AGCCATAAAG 1861 CAAGAGTGTT GGCTGAGGCA ATGAGCCAAG CAAACAGTAA CATACTAGTG CAGAGAAGCA 1921 ATTTTAAAGG CTCTAACAGA ATTATTAAAT GTTTCAACTG TGGCAAAGTA GGGCACATAG 1981 CCAGAAATTG CAGGGCCCCT AGGAAAAAGG GCTGTTGGAA ATGTGGACAG GAAGGACACC 2041 AAATGAAAGA CTGTACTGAG AGGCAGGCTA ATTTTTTAGG GAAAATTTGG CCTTCCCACA 2101 AGGGGAGGCC AGGGAATTTC CTCCAGAACA GACCAGAGCC AACAGCCCCA CCAGCAGAAC 2161 CAACAGCCCC ACCAGCAGAG AGCTTCAGGT TCGAGGAGAC AACCCCCGTG CCGAGGAAGG 2221 AGAAAGAGAG GGAACCTTTA ACTTCCCTCA AATCACTCTT TGGCAGCGAC CCCTTGTCTC 2281 AATAAAAGTA GAGGGCCAGA TAAAGGAGGC TCTCTTAGAC ACAGGAGCAG ATGATACAGT 2341 ATTAGAAGAA ATAGATTTGC CAGGGAAATG GAAACCAAAA ATGATAGGGG GAATTGGAGG 2401 TTTTATCAAA GTAAGACAGT ATGATCAAAT ACTTATAGAA ATTTGTGGAA AAAAGGCTAT 2461 AGGTACAGTA TTAGTAGGGC CTACACCAGT CAACATAATT GGAAGAAATC TGTTAACTCA 2521 GCTTGGATGC ACACTAAATT TTCCAATTAG TCCTATTGAA ACTGTACCAG TAAAATTAAA 2581 ACCAGGAATG GATGGCCCAA AGGTCAAACA ATGGCCATTG ACAGAAGAAA AAATAAAAGC 2641 ATTAACAGCA ATTTGTGAGG AAATGGAGAA GGAAGGAAAA ATTACAAAAA TTGGGCCTGA 2701 TAATCCATAT AACACTCCAG TATTTGCCAT AAAAAAGAAG GACAGTACTA AGTGGAGAAA 2761 ATTAGTAGAT TTCAGGGAAC TCAATAAAAG AACTCAAGAC TTTTGGGAAG TTCAATTAGG 2821 AATACCACAC CCAGCAGGAT TAAAAAAGAA AAAATCAGTG ACAGTGCTAG ATGTGGGGGA 2881 TGCATATTTT TCAGTTCCTT TAGATGAAAG CTTCAGGAAA TATACTGCAT TCACCATACC

FIG. 11A



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2941	TAGTATAAAC	AATGAAACAC	CAGGGATTAG	ATATCAATAT	AATGTGCTGC	CACAGGGATG
3001	GAAAGGAICA	CCAGCAATAT	TCCAGAGTAG	מממסמסמסמר CATG	. איירייייא מאמיי	1 CCM
3001	MANAMATCCA	GACATAGTTA	. TCTATCAATA	ТАТССАТСАС	י איייטייט איייטייטיעיי	
3121	AGMANTAGGG	CAACATAGAG	CAAAAATAGA	. AGAGTTAAGG	CAACA COMMONAC	
3181	ATTTACAACA	CCAGACAAGA	AACATCAAAA	AGAACCCCCA	TTTCTTTCA	TGGGGTATGA
3241	ACTCCATCCT	GACAAATGGA	CAGTACAACC	TATACTCCCA	CCACAAAAA	ATAGTTGGAC
3301	TGTCAATGAT	ATACAGAAGT	TAGTGGGAAA	בבותבוטבוטבוטב בשתי או או או היים או	CCAGAAAAGG	TTTACCCAGG
3361	GATTAAAGTA	AGGCAACTCT	GTAAACTCCT	Cleggeeece	A A A COA COA	CAGACATAGT
3421	ACCACTAACT	GAAGAAGCAG	AATTAGAATT	GGCAGAGAA	AAAGCACTAA	CAGACATAGT TAAGAGAACC
3481	AGTACATGGA	GTATATTATG	ATCCATCAAA	AGA CTTCATA	AGGGAAATTTT	TAAGAGAACC AGAAACAGGG
3541	GCATGAACAA	TGGACATATC	אייים מיייים מממ	ACA ACCAMMO	GCIGAAATAC	AGAAACAGGG AAACAGGGAA
3601	GTATGCAAAA	ATGAGGACTA	CCCACACTAA	TCATCTATA	AAAAATCTGA	AAACAGGGAA AGGCAGTGCA
3661	AAAAATAGCC	ATGGAAAGCA	TAGTAATATG	CCCAAACAC	CAGTTAACAG	AGGCAGTGCA GACTACCCAT
3721	CCAAAAAGAA	ACATGGGAGA	CATGGTGGAG	A CA CER EDGG	CCTAAATTTA	GACTACCCAT GGATCCCTGA
3781	GTGGGAGTTT	GTTAATACCC	CTCCCCTACT	AGACTATTGG	CAAGCCACCT	GGATCCCTGA AAAAAGATCC
3841	CATAGCAGGA	GTAGAAACTT	TOTATOTAGI	MAAATTATGG	TACCAACTAG	AAAAAGATCC CTAAAATAGG
3901	AAAAGCAGGG	TATGTTACTG	ACAGAGGA	TGGAGCAACT	AATAGGGAAG	CTAAAATAGG
3961	AAATCAGAAG	ACTGAGTTAC	ACAGAGGAAG	GCAGAAAATT	GTTACTCTAA	CTAACACAAC
4021	AAACATAGTA	ACAGACTCAC	AGCAATICA	GCTAGCTCTG	CAGGATTCAG	GATCAGAAGT
4081	AAACATAGTA TGACTCAGAG	ATATTTAACC	AGIAIGCATT	AGGAATCATT	CAAGCACAAC	CAGATAAGAG
4141	GTCATGGGTA	CCACCACATA	AAATAATAGA	ACAGTTAATA	AACAAGGAAA	GAATCTACCT
4201	GTCATGGGTA	ACCA A A CTCT	MAGGAATTGG	GGGAAATGAA	CAAGTAGATA	AATTAGTAAG
4261	TAAGGGAATT	AGGAAAGIGI	TGTTTCTAGA	TGGAATAGAT	AAAGCTCAAG	AAGAGCATGA
4321	AAGGTACCAC	GTA CCTA CCT	GAGCAATGGC	TAATGAGTTT	AATCTGCCAC	CCATAGTAGC
4381	AAAAGAAATA	ACTCCACCCA	GIGATAAATG	TCAGCTAAAA	GGGGAAGCCA	TACATGGACA
4441	AGTCGACTGT	GTCCATCTA C	TATGGCAATT	AGATTGTACC	CATTTAGAGG	GAAAAATCAT
4501	CCTGGTAGCA	ACACCATGTAG	CTAGTGGCTA	CATGGAAGCA	GAGGTTATCC	CAGCAGAAAC
4561	AGGACAAGAA	ACAGCATATT	TTATATTAAA	ATTAGCAGGA	AGATGGCCAG	TCAAAGTAAT
4621	ACATACAGAC	CACCAATTO	ATTTTACCAG	TACTGCAGTT	AAGGCAGCCT	GTTGGTGGGC
4681	AGGTATCCAA	CAGGAATITG	GAATTCCCTA	CAATCCCCAA	AGTCAGGGAG	TGGTAGAATC
4741	CATGAATAAA	CAATTAAAGA	AAATAATAGG	ACAAGTAAGA	GATCAAGCTG	AGCACCTTAA
4801	GACAGCAGTA	CCCCAAAIGGCAG	TATTCATTCA	CAATTTTAAA	AGAAAAGGGG	GAATTGGGGG
4861	GTACAGTGCA	ATTATA	TAATAGACAT	AATAGCAACA	GACATACAAA	CTAAAGAATT
4921	ACAAAAACAA	ATTATAAGAA	TICAAAATTT	TCGGGTTTAT	TACAGAGACA	GCAGAGACCC
4001	TATTTGGAAA	GGACCAGCCG	AACTACTCTG	GAAAGGTGAA	GGGGTAGTAG	TAATAGAAGA
5041	TAAAGGTGAC	ATAAAGGTAG	TACCAAGGAG	GAAAGCAAAA	ATCATTAGAG	ATTATGGAAA
5101	ACAGATGGCA	GGTGCTGATT	GTGTGGCAGG	TGGACAGGAT	GAAGATTAGA	GCATGGAATA
5161	GTTTAGTAAA	GCACCATATG	TATATATCAA	GGAGAGCTAG	TGGATGGGTC	TACAGACATC
2101	ATTTTGAAAG	CAGACATCCA	AAAGTAAGTT	CAGAAGTACA	TATCCCATTA	GGGGATGCTA
5221	GATTAGTAAT	AAAAACATAT	TGGGGTTTGC	AGACAGGAGA	AAGAGATTGG	CATTTGGGTC
J201	AIGGAGICIC	CATAGAATGG	AGACTGAGAG	AATACAGCAC	ACAAGTAGAG	CCTCACCTCC
224T	CAGACCAGCT	AATTCACATG	CATTATTTTG	ATTGTTTTAC	AGAATCTCCC	70777777777
2401	CCATATTAGG	ACACATAGTT	TTTCCTAGGT	GTGACTATCA	AGCAGGACAT	AACAACCOBAC
240T	GAICICIGCA	ATACTTGGCA	CTGACAGCAT	TGATAAAACC	AAAAAAAAAA	A A C C C A C C C C C
J J Z I	ICCLIMITAL	TAGAAAATTA	GTAGAGGATA	GATGGAACGA	CCCCCAGAAG	ACCACCCCC
220I	GCAGAGGGAA	CCATACAATG	AATGGACACT	AGAGATTCTA	GAAGAACTCA	ACCACCA ACC
2041	IGICAGACAC	TTTCCTAGAC	CATGGCTCCA	TAGCTTAGGA	CD D TD 	ATCA BACCOS
3/01	IGGGGATACT	TGGACGGGAG	TTGAAGCTAT	AATAAGAGTA	CTGCDDCDDC	TA CTCTTCA m
J / 0 I	ICATITCAGA	ATTGGATGCC	AACATAGCAG	AATAGGCATC	TTCCCACACA	033030033
JU21	MANIGGAGCC	AGTAGATCCT	AAACTAAAGC	CCTGGDDCCD	TCCACCAACC	03300m33
5881	CAGCTTGTAA	TAATTGCTTT	TGCAAACACT	GTAGCTATCA	TTGTCTAGTT	TGCTTTCAGA
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FIG. 11B



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5941	CAAAAGGTTT	AGGCATTTCC	TATGGCAGGA	AGAAGCGGAG	ACAGCGACGA	AGCGCTCCTC
6001	CAAGTGGTGA	AGATCATCAA	AATCCTCTAT	CAAAGCAGTA	AGTACACATA	GTAGATOTA A
909T	TGGTAAGTTT	AAGTTTATTT	' AAAGGAGTAG	ATTATAGATT	AGGAGTAGGA	GCATTCATAC
6121	TAGCACTAAT	CATAGCAATA	ATAGTGTGGA	CCATAGCATA	TATAGAATAT	AGGAAATTGG
6181	TAAGACAAAA	GAAAATAGAC	TGGTTAATTA	AAAGAATTAG	GGAAAGAGCA	GAAGAGAGTG
6241	GCAATGAGAG	TGATGGGGAC	ACAGAAGAAT	TGTCAACAAT	GGTGGATATC	GGGCATCTTA
6301	GGCTTCTGGA	TGCTAATGAT	TTGTAACACG	GAGGACTTGT	GGGTCACAGT	CTACTATGGG
6361	GTACCTGTGT	GGAGAGAAGC	AAAAACTACT	CTATTCTGTG	CATCAGATGG	TAAAGGATAT
6421	GAGACAGAAG	TGCATAATGT	CTGGGCTACA	CATGCTTGTG	TACCCACAGA	CCCCAACCCA
6481	CAAGAAATAG	TTTTGGGAAA	TGTAACAGAA	AATTTTAATA	TGTGGAAAA	TAACAMCCCA
6541	GATCAGATGC	ATGAGGATAT	AATCAGTTTA	TGGGATCAAA	GCCTAAAGCC	ATCTCTA
6601	TTGACCCCAC	TCTGTGTCAC	TTTAAACTGT	ACAGATACAA	ATCTTACACC	MIGIGIAAAG
6661	GTTACAGGTA	ATACAAATGA	TACCAATATT	GCAAATGCTA	CATATAACAGG	TAATAGAACT
6721	AAAAATTGCT	CTTTCAATGC	AACCACAGAA	TTAAGAGATA	ACANACAMAN	1GAAGAAATG
6781	CTCTTTTATA	AACTTGATAT	AGTACCACTT	DATGADATA	CTA A CA A COM	AGAGTATGCA
6841	TTAATAAATT	GCAATACCTC	AACCATAACA	CARCCCTCTC	GIAACAACTT	TACATATAGA
6901	ATTCCTATAC	ATTACTGTGC	TCCAGCTGAT	TATGCGATTC	TARAGGTCTC	TTTTGACCCG
6961	TTCAATGGGA	CAGGACCATG	TTDTDDTCTC	ACCACACTAC	TAAAGTGTAA	TAATAAGACA
7021	CCAGTGGTAT	CAACTCAACT	DCTCTTN N N T	AGCACAGIAC	AATGTACACA	TGGAATTAAG
7081	AGATCTGAAA	ATTTGACAGA	CANTACCANA	ACARDADA	CAGAAGAAGG	GATAATAATT
7141	GAGATTAATT	GTACAAGGCC	CAACAATACCAAA	ACANTANTAG	TACATCTTAA	TGAATCTGTA
7201	CAAGCATTCT	ATGCAACAAA	TCACCATIANT	CCAAGGAAAA	GTGTAAGGAT	AGGACCAGGA
7261	AGTACAGATA	GATGGAATAA	A A COTTON CA A	GGAAACATAA	GACAAGCACA	TTGTAACATT
7321	ССТААТААА	CALCOARIA	TCAACCACAC	CAGGTAATGA	AAAAATTAGG	AGAGCATTTC
7381	AGCTTTAATT	GTAGAGGAGA	ATTTTTTTTTT	GCAGGAGGGG	ATCTAGAAAT	TACAATGCAT
7441	AGCTTTAATT TACTACCCTA	AGAATGGTAC	ATTITICIAT	TGCAATACAT	CAAACCTGTT	TAATAGTACA
7501	TACTACCCTA CAATGCAAAA	TORATOGIAC	TOTACOCAMO	AATGGTAATT	CAAGCTTACC	CATCACACTC
7561	CCTCCCATTG	CAGGAAACAT	AACATCTACA	TGGCAAGGGG	TAGGACAAGC	AATGTATGCC
7621	CCTCCCATTG	TTAACAACAC	AACAIGIAGA	TCAAACATCA	CAGGAATACT	ATTGACACGT
7681	GATGGGGGAT	ATAACTACAC	A A COUCA A COUR	ACAGAGGAGA	CATTCAGACC	TGGAGGAGGA
7741	GATATGAGGG TTGGGAATAG	CACCCACTAA	CCCAAAAA	TATAAATATA	AAGTGGTAGA	AATTAAGCCA
7801	TTGGGAATAG	CTCTCTTCCT	TCCCTTCTTC	AGAGTGGTGC	AGAGAAAAA	AAGAGCAGTG
7861	GGAATAGGAG TCAATAACGC	TGACGGTACA	CCCCACACA	GGAGCAGCAG	GAAGCACTAT	GGGCGCAGCG
7921	TCAATAACGC	ACCCTATACA	GGCCAGACAA	CTGTTGTCTG	GTATAGTGCA	ACAGCAAAGC
7981	AATTTGCTGA	ACCCCACACA	GGCGCAACAG	CATATGTTGC	AACTCACAGT	CTGGGGCATT
8041	AAGCAGCTCC	AGGCGAGAGT	CCTGGCTATA	GAAAGATACC	TAAAGGATCA	ACAGCTCCTA
8101	GGGATTTGGG	AAMGMGAAGG	AAGACTCATC	TGCACCACTG	CTGTGCCTTG	GAACTCCAGT
8161	TGGAGTAATA	AATCTGAAGC	AGATATTTGG	GATAACATGA	CTTGGATGCA	GTGGGATAGA
0101	GAAATTAATA	ATTACACAGA	AACAATATTC	AGGTTGCTTG	AAGACTCGCA	AAACCAGCAG
0221	GAAAAGAATG	AAAAAGATTT	ATTAGAATTG	GACAAGTGGA	ATAATCTGTG	GAATTGGTTT
0201	GACATATCAA	ACTGGCTGTG	GTATATAAAA	ATATTCATAA	TGATAGTAGG	AGGCTTGATA
0.441	GGTTTAAGAA	TAATTTTTGC	TGTGCTCTCT	ATAGTGAATA	GAGTTAGGCA	GGGATACTCA
8401	CCTTTGTCAT	TTCAGACCCT	TACCCCAAGC	CCGAGGGGAC	TCGACAGGCT	CGGAGGAATC
8461	GAAGAAGAAG	GTGGAGAGCA	AGACAGAGAC	AGATCCATAC	GATTGGTGAG	CGGATTCTTG
8521	TCGCTTGCCT	GGGACGATCT	GCGGAGCCTG	TGCCTCTTCA	GCTACCACCG	CTTGAGAGAC
8281	TTCATATTAA	TTGCAGTGAG	GGCAGTGGAA	CTTCTGGGAC	ACAGCAGTCT	CAGGGGACTA
8641	CAGAGGGGGT	GGGAGATCCT	TAAGTATCTG	GGAAGTCTTG	TGCAGTATTG	GGGTCTAGAG
8,01	CTAAAAAAAGA	GTGCTATTAG	TCCGCTTGAT	ACCATAGCAA	TAGCAGTAGC	TGAAGGAAGA
8/61	GATAGGATTA	TAGAATTGGT	ACAAAGAATT	TGTAGAGCTA	TCCTCAACAT	ACCTAGGAGA
8821	ATAAGACAGG	GCTTTGAAGC	AGCTTTGCTA	TAAAATGGGA	GĠCAAGTGGT	CAAAACGCAG
8881	CATAGTTGGA	TGGCCTGCAG	TAAGAGAAAG	AATGAGAAGA	ACTGAGCCAG	CAGCAGAGGG
8941	AGTAGGAGCA	GCGTCTCAAG	ACTTAGATAG	ACATGGGGCA	CTTACAAGCA	GCAACACACC



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9001	TGCTACTAAT	GAAGCTTGTG	CCTGGCTGCA	AGCACAAGAG	GAGGACGGAG	ATGTAGGCTT
9061	TCCAGTCAGA	CCTCAGGTAC	CTTTAAGACC	AATGACTTAT	AAGAGTGCAG	TAGATCTCAG
9121	CTTCTTTTTA	AAAGAAAAGG	GGGGACTGGA	AGGGTTAATT	TACTCTAGGA	AAAGGCAAGA
9181	AATCCTTGAT	TTGTGGGTCT	ATAACACACA	AGGCTTCTTC	CCTGATTGGC	AAAACTACAC
9241	ATCGGGGCCA	GGGGTCCGAT	TCCCACTGAC	CTTTGGATGG	TGCTTCAAGC	TAGTACCAGT
9301	TGACCCAAGG	GAGGTGAAAG	AGGCCAATGA	AGGAGAAGAC	AACTGTTTGC	TACACCCTAT
9361	GAGCCAACAT	GGAGCAGAGG	ATGAAGATAG	AGAAGTATTA	AAGTGGAAGT	TTGACAGCCT
9421	TCTAGCACAC	AGACACATGG	CCCGCGAGCT	ACATCCGGAG	TATTACAAAG	ACTGCTGACA
9481	CAGAAGGGAC	TTTCCGCCTG	GGACTTTCCA	CTGGGGCGTT	CCGGGAGGTG	TGGTCTGGGC
9541	GGGACTTGGG	AGTGGTCACC	CTCAGATGCT	GCATATAAGC	AGCTGCTTTT	CGCTTGTACT
9601	GGGTCTCTCT	CGGTAGACCA	GATCTGAGCC	TGGGAGCTCT	CTGGCTATCT	AGGGAACCCA
9661	CTGCTTAGGC	CTCAATAAAG	CTTGCCTTGA	GTGCTCTAAG	TAGTGTGTGC	CCATCTGTTG
9721	TGTGACTCTG	GTAACTAGAG	ATCCCTCAGA	CCCTTTGTGG	TAGTGTGGAA	AATCTCTAGC
9781	A					

FIG. 11D



15/23

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MAY 1 2 2003

TECH CENTER 1600/2900

SEQ ID NO:34



16/23

975Pol wt until 6aa Int: (SEQ ID NO:35)

TTTTTTAGGGAAGATTTGGCCTTCCCACAAGGGAAGGCCAGGGAATTTCCTTCAGAA CAGAACAGACCCACCACCAGCAGAGAGCTTCAAGTTCGAGGAGACAACCC CCGCTCCGAAGCAGGAGCCGAAAGACAGGGAACCCTTAATTTCCCTCAAATCACTCT TTGGCAGCGACCCCTTGTCTCAATAAAAGTAGGGGGTCAAATAAAGGAGGCTCTCTT AGACACAGGAGCTGATGATACAGTATTAGAAGAAATGAGTTTGCCAGGAAAATGGA AACCAAAAATGATAGGAGGAATTGGAGGTTTTATCAAAGTAAGACAGTATGATCAA ATACTTATAGAAATTTGTGGAAAAAAGGCTATAGGTACAGTATTAATAGGACCTACA CCTGTCAACATAATTGGAAGGAATATGTTGACTCAGCTTGGATGCACACTAAATTTT AAGGTTAAACAATGGCCATTGACAGAAGAGAAAATAAAAGCATTAACAGCAATTTG TGAAGAAATGGAGAAAGGAAAAATTACAAAAATTGGGCCTGAAAATCCATATA ACACTCCAGTATTTGCCATAAAAAAGAAGGACAGTACTAAGTGGAGAAAGTTAGTA GATTTCAGGGAACTTAATAAAAGAACTCAAGACTTTTGGGAAGTTCAATTAGGAATA CCACACCCAGCAGGGTTAAAAAAAAAAAAAATCAGTGACAGTACTGGATGTGGGGGA TGCATATTTTTCAGTTCCTTTAGATGAGGACTTCAGGAAATATACTGCATTCACCATA CCTAGTATAAACAATGAAACACCAGGGATTAGATATCAATATAATGTGCTTCCACAG GGATGGAAAGGATCACCATCAATATTCCAGAGTAGCATGACAAAAATCTTAGAGCC CTTTAGAGCAAGAAATCCAGAAATAGTCATCTATCAATATATGGATGACTTGTATGT AGGATCTGACTTAGAAATAGGGCAACATAGAGCAAAAATAGAGGAGTTAAGAAAAC TTTCTTTGGATGGGTATGAACTCCATCCTGACAAATGGACAGTACAGCCTATAGAG TTGCCAGAAAAGGAAAGCTGGACTGTCAATGATATACAGAAGTTAGTGGGAAAATT AAATTGGGCCAGTCAGATTTACCCAGGAATTAAAGTAAGGCAACTTTGTAAACTCCT TAGGGGGCCAAAGCACTAACAGATATAGTACCACTAACTGAAGAAGCAGAATTAG AATTGGCAGAGAACAGGGAAATTCTAAGAGAACCAGTACATGGAGTATATTATGAC CCATCAAAAGACTTGGTAGCTGAAATACAGAAACAGGGGCATGACCAATGGACATA TCAAATTTACCAAGAACCATTCAAAAACCTGAAAACAGGGAAGTATGCAAAAATGA GGACTGCCCACACTAATGATGTAAAACAGTTAACAGAGGCAGTGCAAAAAATAGCT ATGGAAAGCATAGTAATATGGGGAAAGACTCCTAAATTTAGACTACCCATCCAAAA AGAAACATGGGAGACATGGTGGACAGACTATTGGCAAGCCACCTGGATTCCTGAGT GGGAGTTTGTTAATACCCCTCCCTTAGTAAAATTATGGTACCAGCTAGAGAAGAAC CCATAATAGGAGCAGAAACTTTCTATGTAGATGGAGCAGCTAATAGGGAAACTAAA ATAGGAAAAGCAGGGTATGTTACTGACAGAGGAAGGCAGAAAATTGTTTCTCTAAC AGGATCAGAAGTAAACATAGTAACAGACTCACAGTATGCATTAGGAATCATTCAAG CACAACCAGATAAGAGTGAATCAGAGTTAGTCAACCAAATAATAGAACAATTAATA AAAAAGGAAAAGGTCTACCTGTCATGGGTACCAGCACATAAAGGAATTGGAGGAAA TGAACAAATAGATAAATTAGTAAGTAAGGGAATCAGGAAAGTGCTGTTTCTAGATG **GAATAGAT**

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TECH CENTER 1600/290

MAY 1,2 2003

SEQ ID NO:36

GGCGGCATCGTGATCTACCAGTACATGGACGACCTGTACGTGGGCAGCGGCG



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MAY 1 2 2003

TECH CENTER 1600/2900

SEQ ID NO: 37

GGIVIYQYMDDLYVGSGG



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12_5/1ZA (SEQ ID NO:45)

TGGAAGGGTTAATTTACTCCAGGAAAAGGCAAGAGATCCTTGATTTATGGGTCTATC ACACACAGGCTACTTCCCTGATTGGCAAAACTACACCCGGGACCAGGGGTCAGA TATCCACTGACCTTTGGATGGTGCTTCAAGCTAGTGCCAGTTGACCCAAGGGAAGTA GAAGAGGCCAACGGAGGAGAAGACAACTGTTTGCTACACCCTATGAGCCAGTATGG AATGGATGATGAACACAAAGAAGTGTTACAGTGGAAGTTTGACAGCAGCCTAGCAC GCAGACACCTGGCCCGCGAGCTACATCCGGATTATTACAAAGACTGCTGACACAGA AGGGACTTTCCGCCTGGGACTTTCCACTGGGCGTTCCAGGGGGAGTGGTCTGGGCG GGACTGGGAGTGGCCAGCCCTCAGATGCTGCATATAAGCAGCGGCTTTTCGCCTGTA CTGGGTCTCTCTAGGTAGACCAGATCCGAGCCTGGGAGCTCTCTGTCTATCTGGGGA ACCCACTGCTTAGGCCTCAATAAAGCTTGCCTTGAGTGCTCTAAGTAGTGTGTGCCC ATCTGTTGTGACTCTGGTAACTCTGGTAACTAGAGATCCCTCAGACCCTTTGTGGT AGTGTGGAAAATCTCTAGCAGTGGCGCCCCGAACAGGGACTTGAAAGCGAAAGTGAG ACCAGAGAAGATCTCTCGACGCAGGACTCGGCTTGCTGAAGTGCACTCGGCAAGAG AGGAGAGAGATGGGTGCGAGAGCGTCAATATTAAGAGGGGGAAAATTAGACAAAT GGGAAAAATTAGGTTACGGCCAGGGGGGAGAAAACACTATATGCTAAAACACCTA GTATGGGCAAGCAGAGAGCTGGAAAGATTTGCAGTTAACCCTGGCCTTTTAGAGAC ATCAGACGGATGTAGAC AAATAATAAAACAGCTACAACCAGCTCTTCAGA CAGGAACAGAGGAAATTAGATCATTATTTAACACAGTAGCAACTCTCTATTGTGTAC ATAAAGGGATAGATGTACGAGACACCAAGGAAGCCTTAGACAAGATAGAGGAGGA ACAAAACAAATGTCAGCAAAAAACACAGCAGGCGGAAGCGGCTGACAAAAAGGTC AGTCAAAATTATCCTATAGTGCAGAACCTCCAAGGGCAAATGGTACACCAGGCCAT ATCACCTAGAACCTTGAATGCATGGGTAAAAGTAATAGAGGAGAAGGCTTTTAGCC CAGAGGTAATACCCATGTTTACAGCATTATCAGAAGGAGCCACCCCACAAGATTTA AACACCATGTTAAATACAGTGGGGGGACATCAAGCAGCCATGCAAATGTTAAAAG ATACCATCAATGAGGAGGCTGCAGAATGGGATAGGTTACATCCAGTACATGCAGGG CCTGTTGCACCAGGCCAGATGAGAGAACCAAGGGGAAGTGACATAGCAGGAACTA CTAGTACCCTTCAAGAACAAATAGCATGGATGACAAGTAACCCACCTATCCCAGTA CAGCCCTGTCAGCATTTTAGACATAAAACAAGGACCAAAGGAACCCTTTAGAGACT ATGTAGACCGGTTCTTCAAAACTTTAAGAGCTGAACAATCTACACAAGAGGTAAAA AATTGGATGACAGACACCTTGTTAGTCCAAAATGCGAACCCAGATTGTAAGACCATT TTAAGAGCATTAGGACCAGGGGCTTCATTAGAAGAAATGATGACAGCATGTCAGGG AGTGGGAGGACCTAGCCACAAAGCAAGAGTTTTGGCTGAGGCAATGAGCCAAGCAA ACAATACAAGTGTAATGATACAGAAAAGCAATTTTAAAGGCCCTAGAAGAGCTGTT AAATGTTTCAACTGTGGCAGGGAAGGGCACATAGCCAGGAATTGCAGGGCCCCTAG GAAAAGGGGCTGTTGGAAATGTGGAAAGGAAGGACACCAAATGAAAGACTGTACT GAGAGGCAGGCTAATTTTTTAGGGAAAATTTGGCCTTCCCACAAGGGGAGGCCAGG GAATTTCCTTCAGAGCAGACCAGAGCCAACAGCCCACCACTAGAACCAACAGCCC CACCAGCAGAGAGCTTCAAGTTCAAGGAGACTCCGAAGCAGGAGCCGAAAGACAG GGAACCTTTAACTTCCCTCAAATCACTCTTTGGCAGCGACCCCTTGTCTCAATAAAA

FIG. 16A



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GTAGCGGGCCAAACAAAGGAGGCTCTTTTAGATACAGGAGCAGATGATACAGTACT AGAAGAAATAAACTTGCCAGGAAAATGGAAACCAAAAATGATAGGAGGAATTGGA GGTTTTATCAAAGTAAGACAGTATGATCAAATACTTATAGAAATTTGTGGAAAAAGG GCTATAGGTACAGTATTAGTAGGACCTACACCTGTCAACATAATTGGAAGAAATCTG TTGACTCAGCTTGGATGCACACTAAATTTTCCAATTAGCCCCATTGAAACTGTACCA GTAAAATTAAAGCCAGGAATGGATGGCCCAAAGGTTAAACAATGGCCATTGACAGA ATTACAAAAATTGGGCCTGAAAATCCATATAACACTCCAGTATTTGCCATAAAGAAG AAGGACAGTACAAAGTGGAGAAAATTAGTAGATTTCAGGGAACTCAATAAAAGAAC TCAAGACTTTTGGGAAGTCCAATTAGGAATACCACACCCAGCAGGGTTAAAAAAGA AAAAATCAGTGACAGTACTGGATGTGGGAGATGCATATTTTTCAGTCCCTTTAGATG AGAGCTTCAGAAAATATACTGCATTCACCATACCTAGTATAAACAATGAAACACCA GGGATTAGATATCAATATATGTTCTTCCACAGGGATGGAAAGGATCACCAGCAA TATTCCAGAGTAGCATGACAAGAATCTTAGAGCCCTTTAGAACACAAAACCCAGAA GTAGTTATCTATCAATATATGGATGACTTATATGTAGGATCTGACTTAGAAATAGGG CAACATAGAGCAAAAATAGAGGAGTTAAGAGGACACCTATTGAAATGGGGATTTAC CACACCAGACAAGAACATCAGAAAGAACCCCCATTTCTTTGGATGGGGTATGAAC TCCATCCTGACAAATGGACAGTACAGCCTATACAGCTGCCAGAAAAGGAGAGCTGG ACTGTCAATGATATACAGAAGTTAGTGGGAAAGTTAAACTGGGCAAGTCAGATTTA CCCAGGGATTAAAGTAAGGCAACTGTGTAAACTCCTTAGGGGAGCCAAAGCACTAA CAGACATAGTGCCACTGACTGAAGAAGCAGAATTAGAATTGGCTGAGAACAGGGA AATTCTAAAAGAACCAGTACATGGAGTATATTATGACCCATCAAAAGATTTAATAG CTGAAATACAGAAACAGGGGAATGACCAATGGACATATCAAATTTACCAAGAACC ATTTAAAAATCTGAGAACAGGAAAGTATGCAAAAATGAGGACTGCCCACACTAATG ATGTGAAACAGTTAGCAGAGGCAGTGCAAAAGATAACCCAGGAAAGCATAGTAATA TGGGGAAAAACTCCTAAATTTAGACTACCCATCCCAAAAGAAACATGGGAGACATG GTGGTCAGACTATTGGCAAGCCACCTGGATTCCTGAGTGGGAGTTTGTCAATACCCC TCCCCTAGTAAAATTGTGGTACCAGCTGGAAAAAGAACCCATAGTAGGGGCAGAAA CTTTCTATGTAGATGGAGCAGCCAATAGGGAAACTAAAATAGGAAAAGCAGGGTAT GTCACTGACAAAGGAAGGCAGAAAGTTGTTTCCTTCACTGAAACAACAAATCAGAA GACTGAATTACAAGCAATTCAGCTAGCTTTGCAGGATTCAGGGCCAGAAGTAAACA TAGTAACAGACTCACAGTATGCATTAGGAATCATTCAAGCACAACCAGATAAGAGT GAATCAGAATTAGTCAGTCAAATAATAGAACAGTTGATAAAAAAGGAAAAAGTCTA CCTATCATGGGTACCAGCACATAAAGGAATTGGAGGAAATGAACAAGTAGACAAAT TAGTAAGTAGTGGAATCAGAAAAGTACTGTTTCTAGATGGAATAGATAAAGCTCAA GAAGAGCATGAAAAATATCACAGCAATTGGAGAGCAATGGCTAGTGAGTTTAATCT GCCACCCATAGTAGCAAAGGAAATAGTAGCCAGCTGTGATAAATGTCAGCTAAAAG GGGAAGCCATGCATGGACAAGTCGACTGTAGTCCAGGAATATGGCAATTAGACTGT ACACATTTAGAAGGAAAAATCATCCTAGTAGCAGTCCATGTAGCCAGTGGCTACAT GGAAGCAGAGGTTATCCCAGCAGAAACAGGACAAGAAACAGCATACTTATACTAA AATTAGCAGGAAGATGGCCAGTCAAAGTAATACATACAGATAATGGCAGTAATTTC ACCAGTACCGCAGTTAAGGCAGCCTGTTGGTGGGCAGATATCCAACGGGAATTTGG AATTCCCTACAATCCCCAAAGTCAAGGAGTAGTAGAATCCATGAATAAAGAATTAA

FIG. 16B



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AGAAAATCATAGGGCAAGTAAGAGATCAAGCTGAGCACCTTAAGACAGCAGTACAA ATGGCAGTATTCATTCACAATTTTAAAAGAAAAGGGGGGGATTGGGGGGTACAGTGC AGGGGAGAGAATAATAGACATAGCATCAGACATACAAACTAAAGAATTACAAA AACAAATTATAAAAATTCAAAATTTTCGGGTTTATTACAGAGACAGCAGAGACCCTA TTTGGAAAGGACCAGCCAAACTACTCTGGAAAGGTGAAGGGGCAGTAGTAATACAA GATAATAGTGATATAAAGGTAGTACCAAGAAGGAAAGCAAAAATCATTAAGGACTA TGGAAAACAGATGGCAGGTGCTGATTGTGTGGCAGGTAGACAGGATGAAGATTAGA CCCATTAGGAGATGCCAGGTTAGTAATAAAAACATATTGGGGTCTGCAGACAGGAG AAAGAGCTTGGCATTTGGGTCACGGAGTCTCCATAGAATGGAGATTGAGAAGATAT AGCACACAAGTAGACCCTGACCTGACAGACCAACTAATTCATATGCATTATTTTGAT TGTTTTGCAGAATCTGCCATAAGGAAAGCCATACTAGGACAGATAGTTAGCCCTAA GTGTGACTATCAAGCAGGACATAACAAGGTAGGATCTCTACAATACTTGGCACTGA CAGCATTGATAAAACCAAAAAAGATAAAGCCACCTCTGCCTAGTGTTAGGAAATTA GTAGAGGATAGATGGAACAAGCCCCAGAAGACCAGGGGCCCGCAGAGGGAACCATA CAATGAATGGACACTAGAGCTTTTAGAAGAACTCAAGCAGGAAGCTGTCAGACACT TTCCTAGACCATGGCTCCATAACTTAGGACAACATATCTATGAAACCTATGGAGATA CTTGGACAGGAGTTGAAGCAATAATAAGAATCCTGCAACAATTACTGTTTATTCATT TCAGGATTGGGTGCCATCATAGCAGAATAGGCATTTTGCGACAGAGAAGAGCAAGA AATGGAGCCAATAGATCCTAACCTAGAACCCTGGAACCATCCAGGAAGTCAGCCTA AAACTGCTTGTAATGGGTGTTACTGTAAACGTTGCAGCTATCATTGTCTAGTTTGCTT TCAGAAAAAGGCTTAGGCATTTACTATGGCAGGAAGAAGCGGAGACAGCGACGAA AATAGTATATGTAATGTTAGATTTAACTGCAAGAATAGATTCTAGATTAGGAATAGG AGCATTGATAGTAGCACTAATCATAGCAATAATAGTGTGGACCATAGTATATAG GAAAGAGCAGAAGACAGTGGCAATGAGAGCGAGGGGGGATACTGAAGAATTATCGA CACTGGTGGATATGGGCATCTTAGGCTTTTGGATGCTAATGATGTGAA GGGCTTGTGGGTCACAGTCTACTACGGGGTACCTGTGGGGAGAGAAGCAAAAACT GGCTACACATGCCTGTGTACCCACAGACCCCAACCCACAAGAAGTGATTTTGGGC AATGTAACAGAAAATTTTAACATGTGGAAAAATGACATGGTGGATCAGATGCAGG AAGATATAATCAGTTTATGGGATCAAAGCCTTAAGCCATGTGTAAAA1TGACCCCA CTCTGTGTCACTTTAAACTGTACAAATGCAACTGTTAACTACAATAATACCTCTAAA GACATGAAAAATTGCTCTTTCTATGTAACCACAGAATTAAGAGATAAGAAAAAGAA AGAAAATGCACTTTTTTATAGACTTGATATAGTACCACTTAATAATAGGAAGAATGG GAATATTAACAACTATAGATTAATAAATTGTAATACCTCAGCCATAACACAAGCCTG TCCAAAAGTCTCGTTTGACCCAATTCCTATACATTATTGTGCTCCAGCTGGTTATGCG CCTCTAAAATGTAATAAGAAATTCAATGGAATAGGACCATGCGATAATGTCAG CACAGTACAATGTACACATGGAATTAAGCCAGTGGTATCAACTCAATTACTGTTAAA TCAAAACAATAATAGTACATCTTAATGAATCTATAGAGATTAAATGTACAAGACC

FIG. 16C



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TGGCAATAATACAAGAAAGAGTGTGAGAATAGGACCAGGACAAGCATTCTATGCA ACAGGAGACATAATAGGAGATATAAGACAAGCACATTGTAACATTAGTAAAAATGA ATGGAATACAACTTTACAAAGGGTAAGTCAAAAATTACAAGAACTCTTCCCTAATA GTACAGGGATAAAATTTGCACCACACTCAGGAGGGGACCTAGAAATTACTACACAT AGCTTTAATTGTGGAGGAGAATTTTTCTATTGCAATACAACAGACCTGTTTAATAGT ACATACAGTAATGGTACATGCACTAATGGTACATGCATGTCTAATAATACAGAGCG CATCACACTCCAATGCAGAATAAAACAAATTATAAACATGTGGCAGGAGGTAGGAC GGACTACTATTAACACGTGATGGAGGAGATAATAATACTGAAACAGAGACATTCAG ACCTGGAGGAGGAGACATGAGGGACAATTGGAGAAGTGAATTATATAAATACAAG GTGGTAGAAATTAAACCATTAGGAGTAGCACCCACTGCTGCAAAAAGGAGAGTGGT GGAGAGAAAAAAGAGCAGTAGGAATAGGAGCTGTGTTCCTTGGGTTCTTGGGAG CAGCAGGAAGCACTATGGGCGCAGCATCAATAACGCTGACGGTACAGGCCAGACAA TTATTGTCTGGTATAGTGCAACAGCAAAGTAATTTGCTGAGGGCTATAGAGGCGCAA CAGCATATGTTGCAACTCACGGTCTGGGGCATTAAGCAGCTCCAGGCAAGAGTCCTG GCTATAGAGAGATACCTACAGGATCAACAGCTCCTAGGACTGTGGGGCTGCTCTGG AAAACTCATCTGCACCACTAATGTGCTTTGGAACTCTAGTTGGAGTAATAAAACTCA AAGTGATATTTGGGATAACATGACCTGGATGCAGTGGGATAGGGAAATTAGTAATT TGAAAAAGATTTACTAGCATTGGACAGGTGGAACAATCTGTGGAATTGGTTTAGCAT AACAAATTGGCTGTGGTATATAAAAATATTCATAATGATAGTAGGAGGCTTGATAG GTTTAAGAATAATTTTTGCTGTGCTCTCTCTAGTAAATAGAGTTAGGCAGGGATACT CACCCTTGTCATTGCAGACCCTTATCCCAAACCCGAGGGGACCCGACAGGCTCGGA GGAATCGAAGAAGAAGGTGGAGAGCAAGACAGCAGCAGATCCATTCGATTAGTGA GCGGATTCTTGACACTTGCCTGGGACGACCTACGAAGCCTGTGCCTCTTCTGCTACC ACCGATTGAGAGACTTCATATTAATTGTAGTGAGAGCAGTGGAACTTCTGGGACAC AGTAGTCTCAGGGGACTGCAGAGGGGGGGGGGGAACCCTTAAGTATTTGGGGAGTCT TGTGCAATATTGGGGTCTAGAGTTAAAAAAAGAGTGCTATTAATCTGCTTGATACTAT AGCAATAGCAGTAGCTGAAGGAACAGATAGGATTCTAGAATTCATACAAAACCTTT GTAGAGGTATCCGCAACGTACCTAGAAGAATAAGACAGGGCTTCGAAGCAGCTTTG CAATAAAATGGGGGCAAGTGGTCAAAAAGCAGTATAATTGGATGGCCTGAAGTAA GAGAAAGAATCAGACGAACTAGGTCAGCAGCAGAGGGAGTAGGATCAGCGTCTCA AGACTTAGAGAAACATGGGGCACTTACAACCAGCAACACAGCCCACAACAATGCTG CTTGCGCCTGGCTGGAAGCGCAAGAGGAGGAAGGAGAAGTAGGCTTTCCAGTCAGA CCTCAGGTACCTTTAAGACCAATGACTTATAAAGCAGCAATAGATCTCAGCTTCTTT TTAAAAGAAAAGGGGGACTGGAAGGGTTAATTTACTCCAAGAAAAGGCAAGAGAT CCTTGATTTGTGGGTTTATAACACACAAGGCTTCTTCCCTGATTGGCAAAACTACAC ACCGGGACCAGGGTCAGATTTCCACTGACCTTTGGATGGTACTTCAAGCTAGAGCC AGTCGATCCAAGGGAAGTAGAAGAGGCCAATGAAGGAGAAAACAACTGTTTACTAC ACCCTATGAGCCAGCATGGAATGGAGGATGAAGACAGAGAAGTATTAAGATGGAAG TTTGACAGTACGCTAGCACGCAGACACATGGCCCGCGAGCTACATCCGGAGTATTAC AAAGACTGCTGACACAGAAGGGACTTTCCGCTGGGACTTTCCACTGGGGCGTTCCAG GAGGTGTGGTCTGGGCGGGACAGGGGGAGTGGTCAGCCCTGAGATGCTGCATATAAG CAGCTGCTTTTCGCCTGTACTGGGTCTCTCTAGGTAGACCAGATCTGAGCCCGGGAG

FIG. 16D



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CTCTCTGGCTATCTAGGGAACCCACTGCTTAAGCCTCAATAAAGCTTGCCTTGAGTG CCTTGAGTAGTGTGTCCCGTCTGTTGTGTGACTCTGGTAACTAGAGATCCCTCAGA CCACTTGTGGTAGTGTGGAAAATCTCTAGCA

FIG. 16E